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EXHIBIT I

An alignment of a DI-FRIL, a Pv-FRIL, and a Yam-FRIL of the present invention.

COMPARISON OF THE AMINO ACID SEQUENCES OF D1-FRIL, Pv-FRIL AND
Ss-FRIL OF THE INVENTION

D1-FRIL: AQSLSFSFTK FDPNQEDLIF QGHATSTNNV LQVTKLDSAG NPVSSSAGRV SEQ ID NO: 2
PvFRIL: AQSLSFNFTK FDLDQKDLIF QGDATSTNNV LQLTKLDSGG NPVGASVGRV SEQ ID NO: 6
YamFRIL: AQSVSFTFTK FDSQKDLMF QGHTISSNV IQLTKLDSNG NPVSTSVGRV SEQ ID NO: 8

D1-FRIL: LYSAPLRLWE DSAVLTSFDT IINFEISTPY TSRIADGLAF FIAPPDSVIS
PvFRIL: LFSAPPHLWE NSMAVSSFET NLTIQISTPH PYAADGFAF FLAPHDTVIP
YamFRIL: LYSAPLRLWE SSTVVSTFET TTFQISTPY TSPPGDGLAF FLAPYDTVIP

D1-FRIL: YHGGFLGLFP NANTLNNSST SENQTTTKAA SSNVVAVEFD TYLNPDYGDP
PvFRIL: PNSWGKFLGL YSNVFRNSPT SENQSFQDVN TDSRVVAVEF DTFPNANIDP
YamFRIL: PNSAGNLLGL FPNLNALRNS TTSKETTIDV NAASNNVAV EFDYTPNDNI

D1-FRIL: NYIHIGIDVN SIRSKVTAKW DWQNGKIATA HISYNSVSKR LSVTSYYAGS
PvFRIL: NYRHIGIDVN SIKSKETARW EWQNGKTATA RISYNSASKK STVTTFYPGM
YamFRIL: GPPRYKHIGI DVNSIRSKAT VAWDWQNGKT ATAHISYNSA SKRLSVTTFY

D1-FRIL: KPATLSYDIE LHTVLPWVR VGLSASTGQD KERNTVHWS FTSSLWTNVA
PvFRIL: EVVALSHQVD LHAELPEWVR VGLSASTGEE KQKNTIISWS FTSSLKNEV
YamFRIL: PGGKAVSLSH DVELTQVLPQ WIRVGFSAST GLEK

D1-FRIL: KKENENKYIT RGVL

PvFRIL: KEPKEDMYIA NVVRSYTWIN DVLSYISNK* MYDALNNNHK YVRCSTCMLF

PvFRIL: MKKK